OIPE

DATE: 12/07/2001 RAW SEQUENCE LISTING PATENT APPLICATION: US/09/998,284 TIME: 12:57:57

Input Set : A:\Sequence Listing.txt Output Set: N:\CRF3\12072001\1998284.raw

3 <110> APPLICANT: POULSEN, et al. 5 <120> TITLE OF INVENTION: COMPOSITION 7 <130> FILE REFERENCE: 674509-2035

- **Does Not Comply** Corrected Diskette Needed
- C--> 9 <140> CURRENT APPLICATION NUMBER: US/09/998,284 C--> 9 <141> CURRENT FILING DATE: 2001-11-30 9 <150> PRIOR APPLICATION NUMBER: PCT/IB00/00829 10 <151> PRIOR FILING DATE: 2000-06-02 12 <150> PRIOR APPLICATION NUMBER: GB 9913050.2 13 <151> PRIOR FILING DATE: 1999-06-04 15 <160> NUMBER OF SEQ ID NOS: 2 17 <170> SOFTWARE: PatentIn version 3.1

## ERRORED SEQUENCES

82 <210> SEQ ID NO: 2 83 <211> LENGTH: 546 84 <212> TYPE: PRT 85 <213> ORGANISM: Chondrus crispus 87 <400> SEQUENCE: 2 89 Met Ala Thr Leu Pro Gln Lys Asp Pro Gly Tyr Ile Val Ile Asp Val 90 1 10 93 Asn Ala Gly Thr Pro Asp Lys Pro Asp Pro Arg Leu Pro Ser Met Lys 2.5 97 Gln Gly Phe Asn Arg Arg Trp Ile Gly Thr Asn Ile Asp Phe Val Tyr 101 Val Val Tyr Thr Pro Gln Gly Ala Cys Thr Ala Leu Asp Arg Ala Met 55 105 Glu Lys Cys Ser Pro Gly Thr Val Arg Ile Val Ser Gly Gly His Cys 70 109 Tyr Glu Asp Phe Val Phe Asp Glu Cys Val Lys Ala Ile Ile Asn Val 90 85 113 Thr Gly Leu Val Glu Ser Gly Tyr Asp Asp Asp Arg Gly Tyr Phe Val 100 105 117 Ser Ser Gly Asp Thr Asn Trp Gly Ser Phe Lys Thr Leu Phe Arg Asp 120 121 His Gly Arg Val Leu Pro Gly Gly Ser Cys Tyr Ser Val Gly Leu Gly 130 135 140 125 Gly His Ile Val Gly Gly Gly Asp Gly Ile Leu Ala Arg Leu His Gly 150 155 129 Leu Pro Val Asp Trp Leu Ser Gly Val Glu Val Val Lys Pro Val 170 165 133 Leu Thr Glu Asp Ser Val Leu Lys Tyr Val His Lys Asp Ser Glu Gly 180 185 137 Asn Asp Gly Glu Leu Phe Trp Ala His Thr Gly Gly Gly Gly Asn 200 141 Phe Gly Ile Ile Thr Lys Tyr Tyr Phe Lys Asp Leu Pro Met Ser Pro RAW SEQUENCE LISTING
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	142		210					215					220				
				Val	Ile	Δla	Ser		T.e.ii	His	Phe	Ser		Asp	Glv	Phe	Thr
		225	GLY	vul		11±u	230	11011	Leu		2	235			011	1	240
			3.00	7.7.	Leu	~1 n		T 011	T 011	mbx	Tyra		Dho	Tarc	T 011	7.1 -	
		Arg	ASP	Ald	ьeu		Asp	Leu	ьeu	1111	250	тут	FIIC	пур	пеп	255	ALG
	150		_	_	_	245	1		<b>~</b> 1	<b>-</b>		<b>a</b> 1	-1-	Dl	77.		
			Asp	Trp	Lys	Asn	Thr	Val	GTĀ		Phe	GIn	IIe	Pne		GIn	Ата
	154				260					265					270		
	157	Ala	Glu		Phe	Val	Met	$\mathtt{Tyr}$	Leu	Tyr	Thr	Ser	Tyr		Asn	Asp	Ala
	158			275					280					285			
	161	Glu	Arg	Glu	Val	Ala	Gln	Asp	Arg	His	Tyr	His	Leu	Glu	Ala	Asp	Ile
	162		290					295					300				
	165	Glu	Gln	Ile	Tyr	Lys	Thr	Cys	Glu	Pro	Thr	Lys	Ala	Leu	Gly	Gly	His
	166	305					310					315			٠		320
	169	Ala	Gly	Trp	Ala	Pro	Phe	Pro	Val	Arg	Pro	Arg	Lys	Arg	His	Thr	Ser
	170		-	-,		325				-	330	_	_	_		335	
		Lvs	Thr	Ser	Tyr	Met	His	Asp	Glu	Thr	Met	asp	Tvr	Pro	Phe	Tyr	Ala
	174	-1-			340					345			- 4 -		350		
		T.e.11	Thr	Glu	Thr	Tle	Asn	Glv	Ser		Pro	Asn	Gln	Arα	-	Lvs	Tvr
	178	пси	1111	355		110		017	360				0111	365	011		-1-
		T 110	Cor		Tyr	Mot	T10	Tvc		Dho	Dro	λcn	Dha		Tla	λen	Val
		гу		АТА	тХт	Met	TIE	375	ASP	FILE	FIO	АБР		GIII	116	АЗР	Vai
	182	-3.	370	•	m	· •	m1		37- 3	D	3	C1	380	m\	C	71-	C1
			Trp	гàг	Tyr	ьeu		GIU	vaı	Pro	Asp	_	ьeu	THE	ser	Ата	
		385		_		_	390	~ 3		_		395	<b>~</b> 3	<b>~</b> 1	<b>61</b>	~1.	400
		Met	Lys	Asp	Ala		Leu	GIn	val	Asp		Pne	GTĀ	GIA	GIU		HIS
	190					405	_				410		_		_	415	
	193	Lys	Val	Val	$\mathtt{Trp}$	Asp	Ala	Thr	Ala		Ala	Gln	Arg	GLu		IIe	IIe
	194				420					425					430		
	197	Lys	Leu	Gln	Tyr	Gln	Thr	${ t Tyr}$	_	Gln	Glu	Glu	Asp		Asp	Ala	Val
	198			435					440					445			
	201	Asn	Leu	Lys	Trp	Ile	Arg	Asp	Phe	Tyr	Glu	Glu	Met	Tyr	Glu	Pro	Tyr
	202		450					455					460				
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•	206	465					470					475					480
	209	Phe	Glu	Gly	Cys	Tyr	Phe	Asn	Tyr	Pro	Asp	Val	Asp	Leu	Asn	Asn	Trp
	210					485					490					495	
	213	Lys	Asn	Gly	Lys	Tyr	Gly	Ala	Leu	Glu	Leu	Tyr	Phe	Leu	Gly	Asn	Leu
	214	-		•	500	-	-			505		_			510		
	217	Asn	Ara	Leu	Ile	Lvs	Ala	Lvs	Trp	Leu	Trp	Asp	Pro	Asn	Glu	Ile	Phe
	218		5	515		_1_		-1-	520		1			525			,
		Thr	Δen		Gln	Ser	Tle	Pro		Lvs	Pro	Len	Lvs		Pro	Lvs	Gln
	222	T111	530	בות	0111	001	110	535	+ - 1 - 1	_13	110	LCu	540	O L U		-15	O 111
		Thr						555			•		540				
			-														
		545/															
	232	(34 )															

VERIFICATION SUMMARY

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Input Set : A:\Sequence Listing.txt
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L:9 M:270 C: Current Application Number differs, Replaced Current Application No

L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:232 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:2